

04/00

Page 1 of 9

SW

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OIPE

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/021,811

DATE: 01/15/2002  
TIME: 18:07:38

Input Set : A:\BB1294 USDIV Seq.txt  
Output Set: N:\CRF3\01152002\J021811.raw

3 <110> APPLICANT: Cahoon, Rebecca E.  
4 Fang, Yiwen  
5 Odell, Joan  
6 Weng, Zude  
8 <120> TITLE OF INVENTION: Plant Myb Transcription Factor Homologs  
10 <130> FILE REFERENCE: BB1294 US NA  
12 <140> CURRENT APPLICATION NUMBER: US/10/021,811  
13 <141> CURRENT FILING DATE: 2001-12-14  
15 <150> PRIOR APPLICATION NUMBER: 60/110,609  
W--> 16 <151> PRIOR FILING DATE: 1998-December-02  
18 <160> NUMBER OF SEQ ID NOS: 63  
20 <170> SOFTWARE: Microsoft Office 97  
22 <210> SEQ ID NO: 1  
23 <211> LENGTH: 771  
24 <212> TYPE: DNA  
25 <213> ORGANISM: Zea mays  
27 <220> FEATURE:  
28 <221> NAME/KEY: unsure  
29 <222> LOCATION: (4) ^  
31 <220> FEATURE:  
32 <221> NAME/KEY: unsure  
33 <222> LOCATION: (99) ^  
35 <220> FEATURE:  
36 <221> NAME/KEY: unsure  
37 <222> LOCATION: (396) ^  
39 <220> FEATURE:  
40 <221> NAME/KEY: unsure ^  
41 <222> LOCATION: (470)  
43 <220> FEATURE:  
44 <221> NAME/KEY: unsure ^  
45 <222> LOCATION: (486)  
47 <220> FEATURE:  
48 <221> NAME/KEY: unsure ^  
49 <222> LOCATION: (586)  
51 <220> FEATURE:  
52 <221> NAME/KEY: unsure ^  
53 <222> LOCATION: (600)  
55 <220> FEATURE:  
56 <221> NAME/KEY: unsure ^  
57 <222> LOCATION: (612)  
59 <220> FEATURE:  
60 <221> NAME/KEY: unsure ^  
61 <222> LOCATION: (615)  
63 <220> FEATURE:  
64 <221> NAME/KEY: unsure ^  
65 <222> LOCATION: (627)  
67 <220> FEATURE:

ENTERED

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/021,811

DATE: 01/15/2002  
TIME: 18:07:38

Input Set : A:\BB1294 USDIV Seq.txt  
Output Set: N:\CRF3\01152002\J021811.raw

68 <221> NAME/KEY: unsure  
 69 <222> LOCATION: (632) /  
 71 <220> FEATURE:  
 72 <221> NAME/KEY: unsure  
 73 <222> LOCATION: (658) /  
 75 <220> FEATURE:  
 76 <221> NAME/KEY: unsure  
 77 <222> LOCATION: (690) /  
 79 <220> FEATURE:  
 80 <221> NAME/KEY: unsure  
 81 <222> LOCATION: (716) /  
 83 <220> FEATURE:  
 84 <221> NAME/KEY: unsure  
 85 <222> LOCATION: (724) /  
 87 <220> FEATURE:  
 88 <221> NAME/KEY: unsure  
 89 <222> LOCATION: (736) /  
 91 <220> FEATURE:  
 92 <221> NAME/KEY: unsure  
 93 <222> LOCATION: (752) /  
 95 <220> FEATURE:  
 96 <221> NAME/KEY: unsure  
 97 <222> LOCATION: (758) /  
 99 <400> SEQUENCE: 1  
 W--> 100 caanacgcggg attgttcaat ccgttcgaca tcacaaaatc cacgcacaaa gaagcgacag 60  
 W--> 101 atgactacga gcagggtggc caggtcggtc ggccgcggna ggcacgtga gccggcggtg 120  
 102 cgcaaggggc cgtggacgt ggaggaggac ctcatcctcg ttagtacat ctcccagcac 180  
 103 ggggaggggct cctgggacaa cctcgcgcgc gcagctggac tgaaccgcaa cggcaagagc 240  
 104 tgcaggctgc ggtggctcaa ctacctgagg cgggggtgc ggcgcggcag catcacggcg 300  
 105 ggggaggaca cggtcatccg ggagctccac gcgagggtggg ggaacaagtg gtccaagatc 360  
 W--> 106 tccaaggcacc tccccggccg aaccgacaaac gagatnaaga actactggag gaccaggatc 420  
 W--> 107 caacaagaag aacagcaagg agccaagacg acgcaacaac gggaccgtcn acgaccgcca 480  
 W--> 108 actccnnggc cgggggacga ctactgggtc cacaacccga ccccgacaaac aagccatact 540  
 W--> 109 gcctgcaaaaa accccatgca actgcacgcg acaacaacccg gtctcntaac aacaagacan 600  
 W--> 110 ccccttcggg gnctnacaac cagaaanccc chccggcggg gaatggtaat cacaacaahaa 660  
 W--> 111 attgtaccct ctgtccaaact aactttcccn cggcacataa acgtcggtc accttnacaa 720  
 W--> 112 tcantcttct ccactnatgc acttgcaac gngtgtantt tgataaacct t 771  
 114 <210> SEQ ID NO: 2  
 115 <211> LENGTH: 157  
 116 <212> TYPE: PRT  
 117 <213> ORGANISM: Zea mays  
 119 <220> FEATURE:  
 120 <221> NAME/KEY: UNSURE  
 121 <222> LOCATION: (111) /  
 123 <220> FEATURE:  
 124 <221> NAME/KEY: UNSURE  
 125 <222> LOCATION: (136) /  
 127 <400> SEQUENCE: 2  
 128 Thr Thr Ser Arg Val Ala Arg Ser Cys Gly Arg Ser Asp Asp Glu

RAW SEQUENCE LISTING  
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Input Set : A:\BB1294 USDIV Seq.txt  
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129 1 5 10 15  
 131 Pro Ala Val Arg Lys Gly Pro Trp Thr Leu Glu Glu Asp Leu Ile Leu  
 132 20 25 30  
 134 Val Ser Tyr Ile Ser Gln His Gly Glu Gly Ser Trp Asp Asn Leu Ala  
 135 35 40 45  
 137 Arg Ala Ala Gly Leu Asn Arg Asn Gly Lys Ser Cys Arg Leu Arg Trp  
 138 50 55 60  
 140 Leu Asn Tyr Leu Arg Pro Gly Val Arg Arg Gly Ser Ile Thr Ala Gly  
 141 65 70 75 80  
 143 Glu Asp Thr Val Ile Arg Glu Leu His Ala Arg Trp Gly Asn Lys Trp  
 144 85 90 95  
 W--> 146 Ser Lys Ile Ser Lys His Leu Pro Gly Arg Thr Asp Asn Glu Xaa Lys  
 147 100 105 110  
 149 Asn Tyr Trp Arg Thr Arg Ile Gln Gln Glu Glu Gln Gln Gly Ala Lys  
 150 115 120 125  
 W--> 152 Thr Thr Gln Gln Arg Asp Arg Xaa Arg Pro Pro Thr Pro Gly Pro Gly  
 153 130 135 140  
 155 Asp Asp Tyr Trp Val His Asn Pro Thr Pro Thr Thr Ser  
 156 145 150 155  
 158 <210> SEQ ID NO: 3  
 159 <211> LENGTH: 782  
 160 <212> TYPE: DNA  
 161 <213> ORGANISM: Zea mays  
 163 <220> FEATURE:  
 164 <221> NAME/KEY: unsure  
 165 <222> LOCATION: (3) ↗  
 167 <400> SEQUENCE: 3  
 wOK> 168 acngtctgct gcaggtacgg gccgtaatcc gggtcgacca cgcgtcccgca caaagtggca 60  
 169 tactttctc tgcgtactgat ttctttttcc ttcctcttc ctcacaaaca gactggattt 120  
 170 caacaagata atcttgaaac tggagccaaac aagcacacag agaaagaaga gcaagaagac 180  
 171 cggctcccgccat cggatatacaag gttaggatgtc gcagcggttag tttcatcata tcgcataaggc 240  
 172 gatatggta cagtggaga ggagactcgc aaggggccat ggacagagaca ggaggacctg 300  
 173 caactggat gcactgtccg tctgttcgtt gaacgtcggt gggatttcata tgccaaagta 360  
 174 tcaggactca accggacagg caagagctgc cggctgcgggt gggtaacta cctccaccct 420  
 175 ggcctcaagc gtgggcgcataatccccat gaagagcgcc tcatccttga gctgcacgc 480  
 176 cgggtggaa acaggtggtc caggatagca cggcgcttgc cagggcgcac tgacaatgag 540  
 177 atcaagaact actggaggac acacatgagg aagaaagcac aggagaggaa gaggaacatg 600  
 178 tctccatcat catccatcat ttcactgagt taccagtca gctacccaga tactccatca 660  
 179 atcattggag ttaaggaca ggagcttcat ggtggcagtgc gtcgcatac aagcatcctg 720  
 180 aagggcaccc atccggacat ggtggctat cccatggacc agatatggat ggaattgaag 780  
 181 gg 782  
 183 <210> SEQ ID NO: 4  
 184 <211> LENGTH: 179  
 185 <212> TYPE: PRT  
 186 <213> ORGANISM: Zea mays  
 188 <400> SEQUENCE: 4  
 189 Met Val Thr Val Arg Glu Glu Thr Arg Lys Gly Pro Trp Thr Glu Gln  
 190 1 5 10 15  
 192 Glu Asp Leu Gln Leu Val Cys Thr Val Arg Leu Phe Gly Glu Arg Arg

RAW SEQUENCE LISTING  
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TIME: 18:07:38

Input Set : A:\BB1294 USDIV Seq.txt  
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193	20	25	30
195	Trp Asp Phe Ile Ala Lys Val Ser Gly Leu Asn Arg Thr Gly Lys Ser		
196	35	40	45
198	Cys Arg Leu Arg Trp Val Asn Tyr Leu His Pro Gly Leu Lys Arg Gly		
199	50	55	60
201	Arg Met Ser Pro His Glu Glu Arg Leu Ile Leu Glu Leu His Ala Arg		
202	65	70	75
204	Trp Gly Asn Arg Trp Ser Arg Ile Ala Arg Arg Leu Pro Gly Arg Thr		
205	85	90	95
207	Asp Asn Glu Ile Lys Asn Tyr Trp Arg Thr His Met Arg Lys Lys Ala		
208	100	105	110
210	Gln Glu Arg Lys Arg Asn Met Ser Pro Ser Ser Ser Ser Ser Leu		
211	115	120	125
213	Ser Tyr Gln Ser Gly Tyr Pro Asp Thr Pro Ser Ile Ile Gly Val Lys		
214	130	135	140
216	Gly Gln Glu Leu His Gly Gly Ser Gly Cys Ile Thr Ser Ile Leu Lys		
217	145	150	155
219	Gly Thr His Pro Asp Met Asp Gly Tyr Pro Met Asp Gln Ile Trp Met		
220	165	170	175
222	Glu Leu Lys		
225	<210> SEQ ID NO: 5		
226	<211> LENGTH: 601		
227	<212> TYPE: DNA		
228	<213> ORGANISM: Zea mays		
230	<220> FEATURE:		
231	<221> NAME/KEY: unsure		
232	<222> LOCATION: (451)		
234	<220> FEATURE:		
235	<221> NAME/KEY: unsure		
236	<222> LOCATION: (456)		
238	<220> FEATURE:		
239	<221> NAME/KEY: unsure		
240	<222> LOCATION: (478)		
242	<220> FEATURE:		
243	<221> NAME/KEY: unsure		
244	<222> LOCATION: (480)		
246	<220> FEATURE:		
247	<221> NAME/KEY: unsure		
248	<222> LOCATION: (490)		
250	<220> FEATURE:		
251	<221> NAME/KEY: unsure		
252	<222> LOCATION: (510)		
254	<220> FEATURE:		
255	<221> NAME/KEY: unsure		
256	<222> LOCATION: (542)		
258	<220> FEATURE:		
259	<221> NAME/KEY: unsure		
260	<222> LOCATION: (549)		
262	<220> FEATURE:		

RAW SEQUENCE LISTING  
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Input Set : A:\BB1294 USDIV Seq.txt  
Output Set: N:\CRF3\01152002\J021811.raw

```

263 <221> NAME/KEY: unsure
264 <222> LOCATION: (552)
266 <220> FEATURE:
267 <221> NAME/KEY: unsure
268 <222> LOCATION: (554) /
270 <220> FEATURE:
271 <221> NAME/KEY: unsure
272 <222> LOCATION: (572) r
274 <220> FEATURE:
275 <221> NAME/KEY: unsure
276 <222> LOCATION: (578) {
278 <220> FEATURE:
279 <221> NAME/KEY: unsure
280 <222> LOCATION: (583) (
282 <220> FEATURE:
283 <221> NAME/KEY: unsure
284 <222> LOCATION: (588) r
286 <220> FEATURE:
287 <221> NAME/KEY: unsure
288 <222> LOCATION: (595)
290 <220> FEATURE:
291 <221> NAME/KEY: unsure
292 <222> LOCATION: (601) r
294 <400> SEQUENCE: 5
295 aaccggccat catcggttat acctaccagc tcgcgtttct tgctgaagcc ctggagctat 60
296 atagcttcga tctgcgcagc acagggttgtc tgtcgactag tgatttagtga agaagatggc 120
297 ggcgcgtgac caccgagagc tgagcggcga cgaggactcc gtgggtggcgg ccggagacct 180
298 cccgcgcggg ccgtggacgg tggaggagga catgtcctc gtcaactacg tcgcgcgcga 240
299 cggcgagggc cgctggaaacg cgctggcacg atgcgcaggg ctccggcgga cgggaaagag 300
300 ctgcgcctg cggtggtctca actacctgcg gccgacactg cggcggggca acatcacggc 360
301 gcaagagcaa ctgctcatcc tggagctgca ctccgcgtgg ggcaaccgct ggtcaagatc 420
W--> 302 ggcgcagcacc tccaaggggca acgacaacga natcanaact actggcgcac cggttcanan 480
W--> 303 caccgcagcan ctcaatgcaa ctcaaagcan cgctcaagga ctcagcgcta atctggatgc 540
W--> 304 gngctccna angnaccgtc gacatccggg angggctnct ttngagcnca cccancaaac 600
W--> 305 n 601
307 <210> SEQ ID NO: 6
308 <211> LENGTH: 120
309 <212> TYPE: PRT
310 <213> ORGANISM: Zea mays
312 <220> FEATURE:
313 <221> NAME/KEY: UNSURE
314 <222> LOCATION: (101)
316 <220> FEATURE:
317 <221> NAME/KEY: UNSURE
318 <222> LOCATION: (113)..(114)
320 <400> SEQUENCE: 6
321 Met Ala Ala Arg Asp His Arg Glu Leu Ser Gly Asp Glu Asp Ser Val
322 1 5 10 15
324 Val Ala Ala Gly Asp Leu Arg Arg Gly Pro Trp Thr Val Glu Glu Asp

```

→ Use of n and/or Xaa has been detected in the Sequence Listing.  
Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY  
PATENT APPLICATION: US/10/021,811

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Input Set : A:\BB1294 USDIV Seq.txt  
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L:12 M:270 C: Current Application Number differs, Replaced Application Number  
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:16 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD  
L:100 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:1  
L:100 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:101 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:1  
L:101 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:106 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:1  
L:106 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:107 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:1  
L:107 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:108 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:1  
L:108 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:109 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:1  
L:109 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:110 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:1  
L:110 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:111 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:1  
L:111 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:112 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:1  
L:112 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:146 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:2  
L:146 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2  
L:152 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:2  
L:152 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2  
L:168 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:3  
L:168 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3  
L:302 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:5  
L:302 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5  
L:303 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:5  
L:303 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5  
L:304 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:5  
L:304 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5  
L:305 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:5  
L:305 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5  
L:339 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:6  
L:339 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6  
L:342 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:6  
L:342 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6  
L:376 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:7  
L:376 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7  
L:377 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:7  
L:377 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7  
L:378 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:7  
L:378 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7  
L:379 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:7  
L:379 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7  
L:402 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:8

VERIFICATION SUMMARY  
PATENT APPLICATION: US/10/021,811

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TIME: 18:07:39

Input Set : A:\BB1294 USDIV Seq.txt  
Output Set: N:\CRF3\01152002\J021811.raw

L:402 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8  
L:638 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:15  
L:638 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15  
L:659 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:16  
L:659 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16  
L:689 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:17  
L:689 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17  
L:825 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:21  
L:825 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21  
L:828 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:21  
L:828 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21  
L:829 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:21  
L:829 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21  
L:830 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:21  
L:830 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21  
L:859 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:22  
L:859 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22  
L:868 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:22  
L:868 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22  
L:941 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:23  
L:941 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23  
L:944 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:23  
L:944 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23  
L:945 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:23  
L:945 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23  
L:946 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:23  
L:946 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23  
L:947 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:23  
L:947 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23  
L:1000 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:24  
L:1000 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24  
L:1012 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:24  
L:1012 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24  
L:1015 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:24  
L:1015 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24  
L:1018 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:24  
L:1018 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24  
L:1021 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:24  
L:1021 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24  
L:1192 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:27  
L:1192 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27  
L:1193 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:27  
L:1193 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27  
L:1194 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:27  
L:1194 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27  
L:1196 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:27  
L:1196 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27  
L:1197 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:27  
L:1197 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/10/021,811

DATE: 01/15/2002

TIME: 18:07:39

Input Set : A:\BB1294 USDIV Seq.txt

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L:1198 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:27  
L:1198 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27  
L:1199 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:27  
L:1199 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27  
L:1222 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:28  
L:1222 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28